

Fig. 1

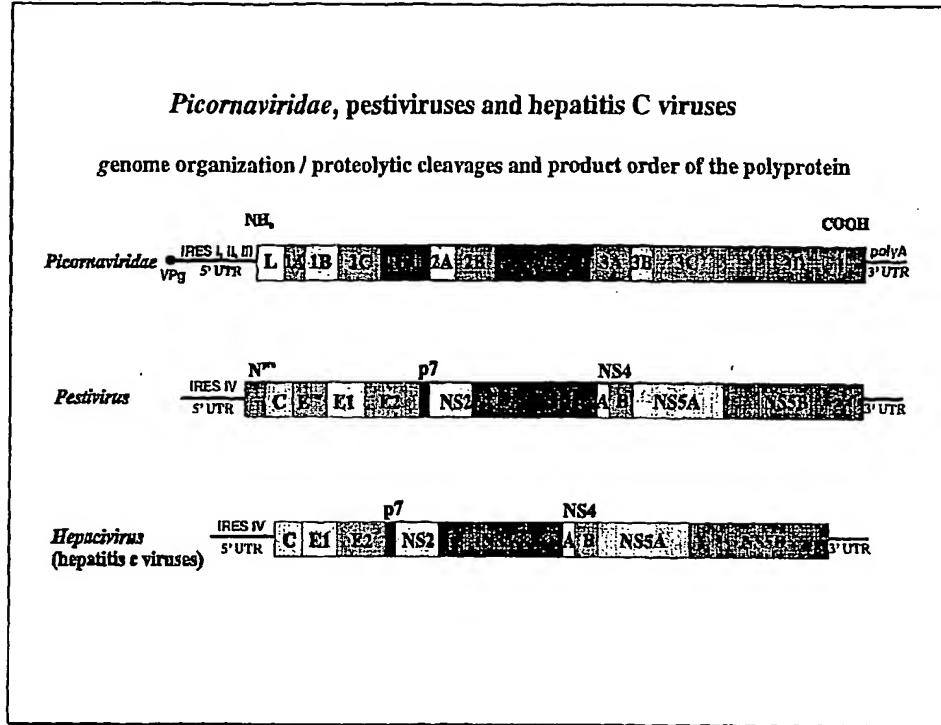


Fig. 1B

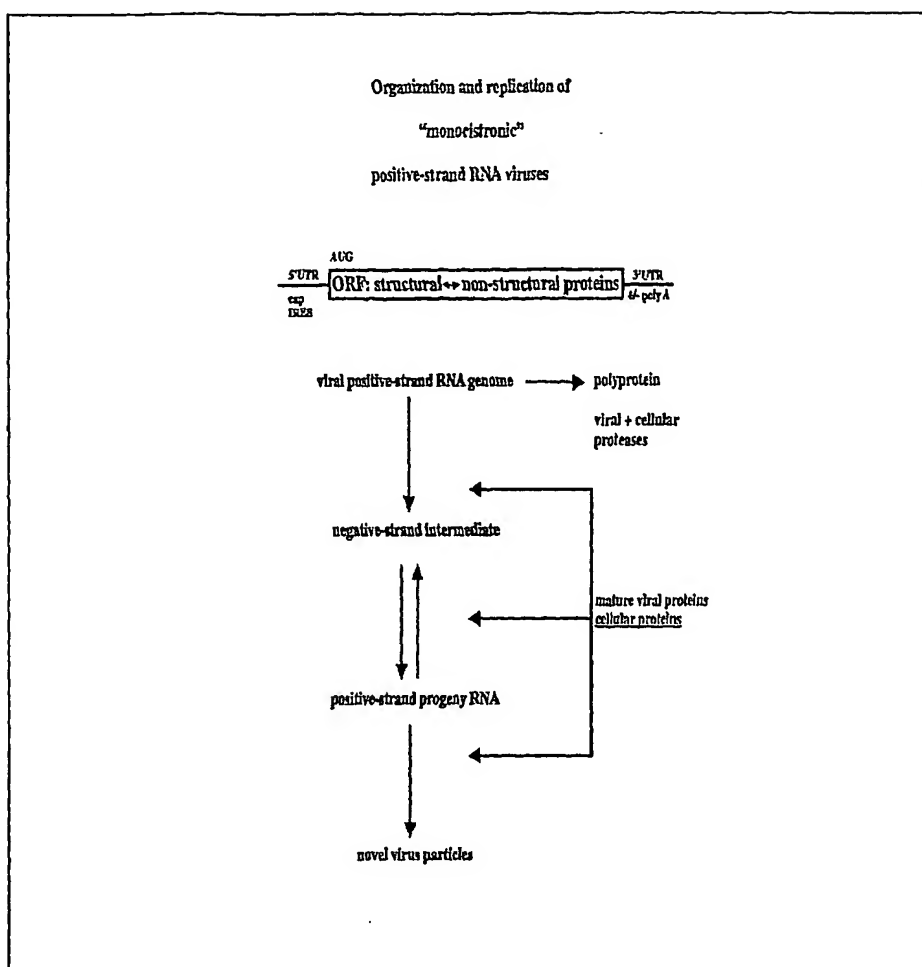


Fig. 2

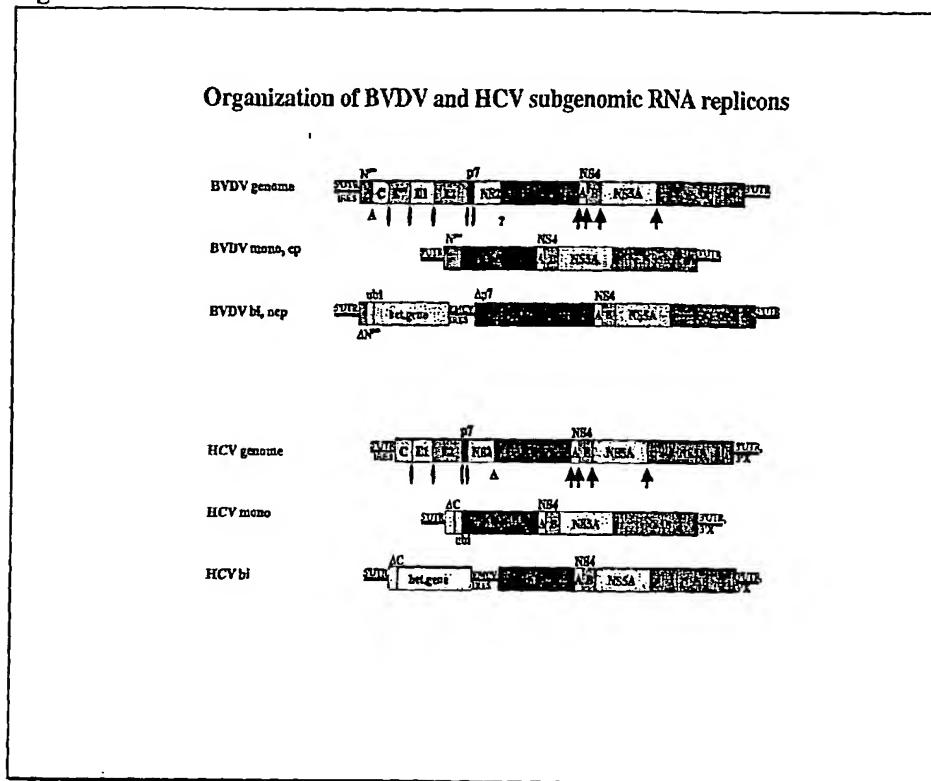


Fig. 3

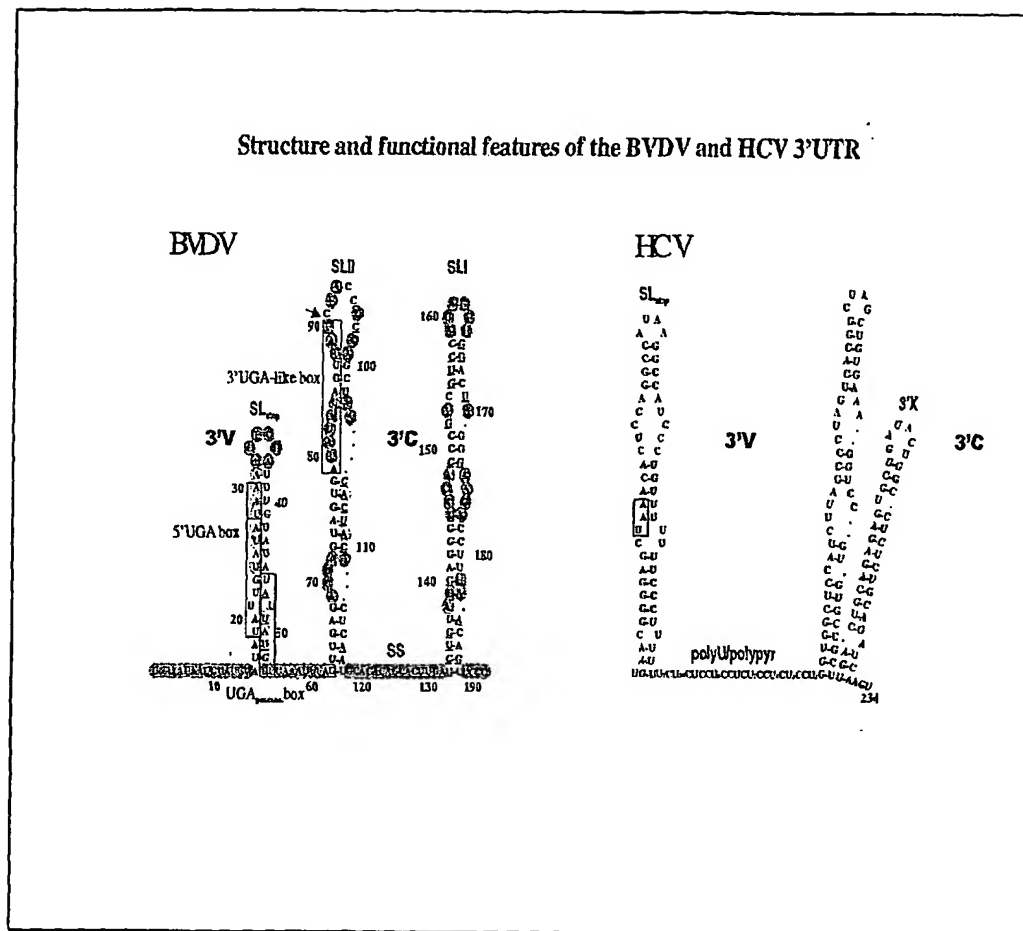


Fig. 4

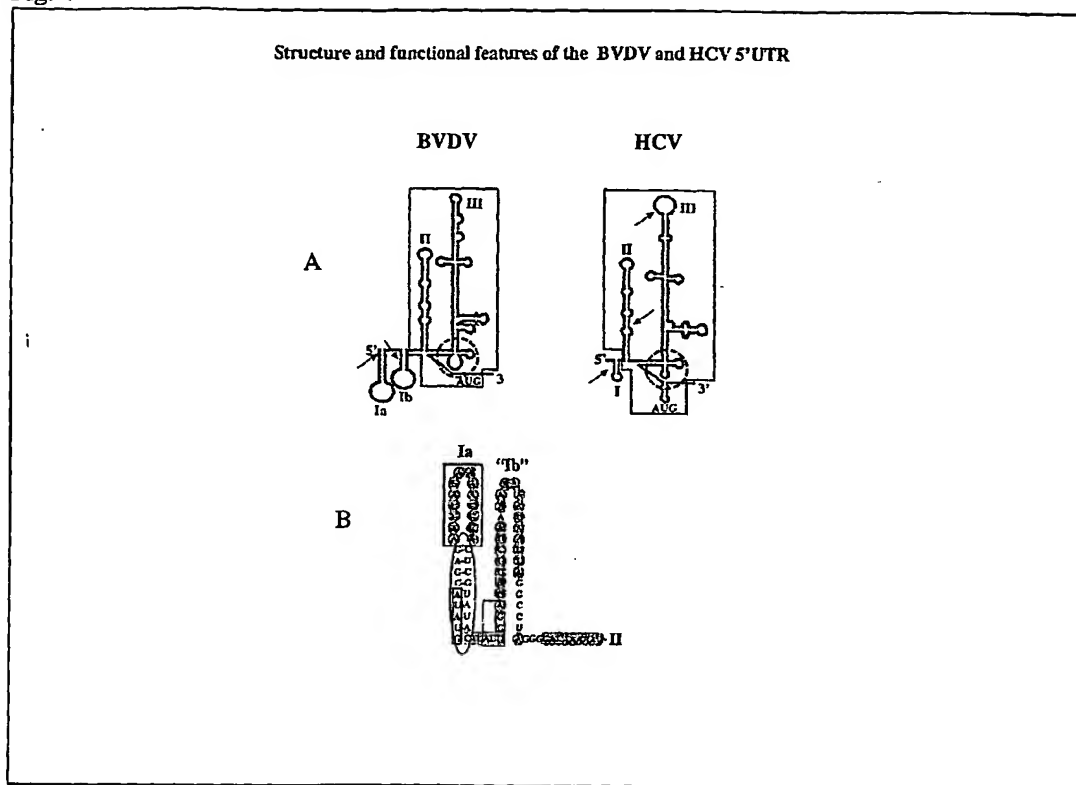


Fig. 5A

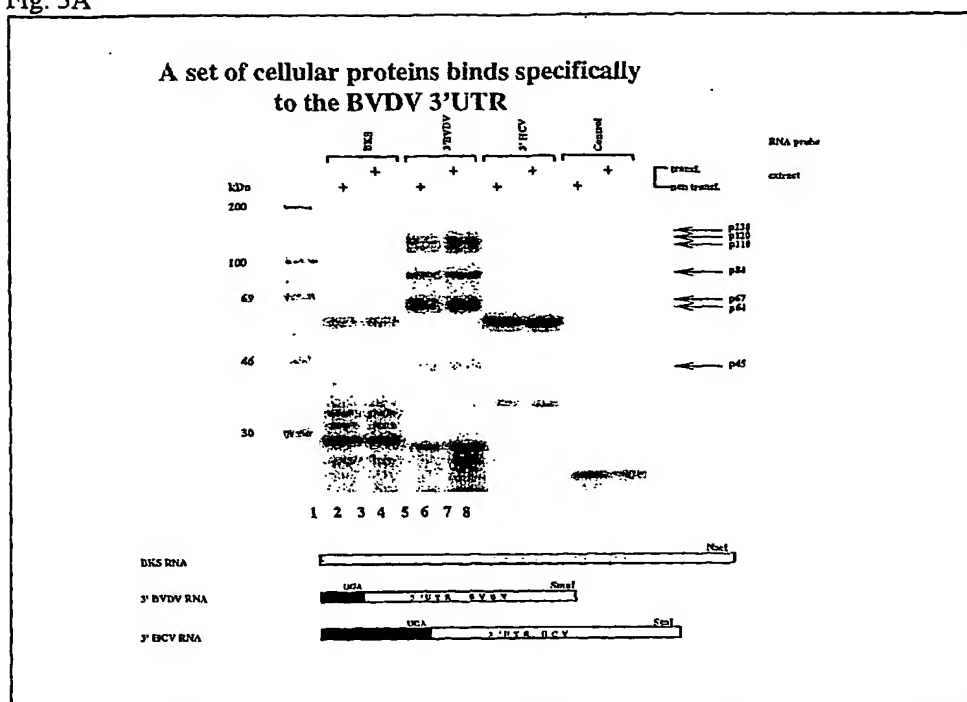


Fig. 5B

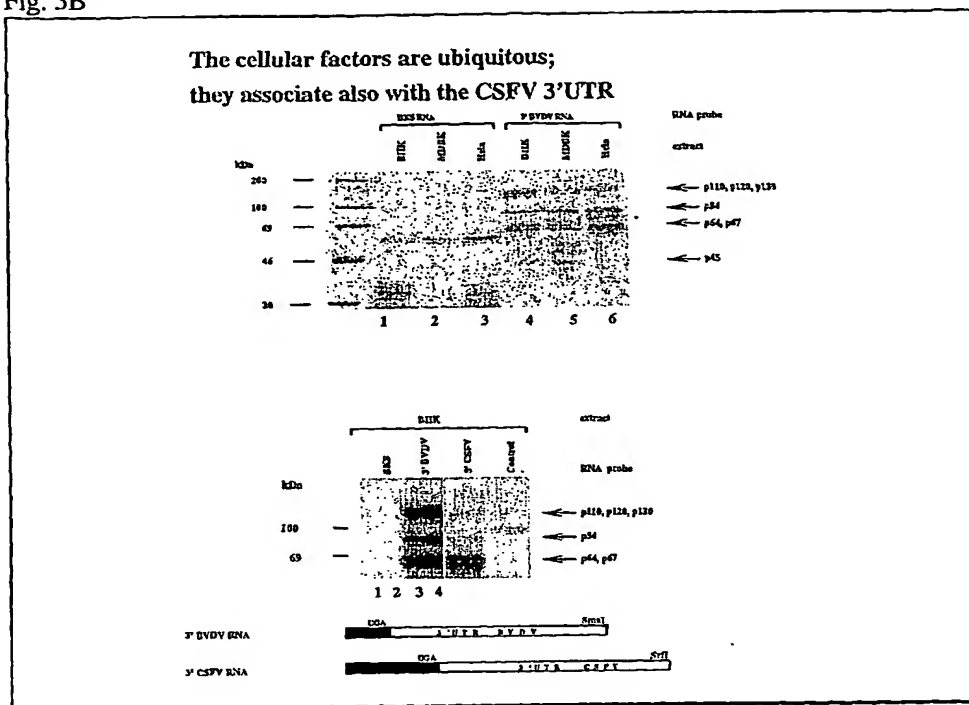


Fig. 5C

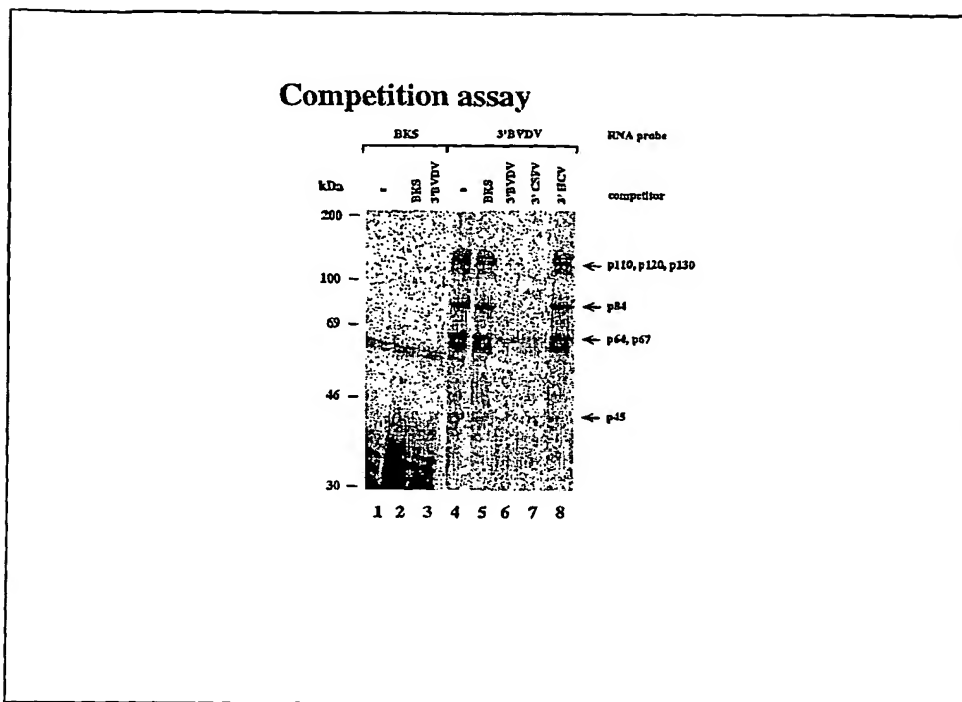


Fig. 5D

Sequence alignment of pestiviral 3'V regions

pestivirus strain	"UGA box" / position
BDV X218	46 AΔU UGU AUA UΔU 57
BDV A841/1	46 GΔU UGU AUA UAA 57
CSFV C-strain	6 UΔU UGU AGA UAA 54
CSFV Alfort-T	43 UΔU UGU AAA UAA 54
BVDV-1 NADL	44 UΔG UGU AUA UAA 55
BVDV-1 Orlow	43 UΔU UAU AUA UAA 54
	16 UΔU UGU ACA UAA 57
BVDV-1 CP7	44 UΔU UAU GUU UAA 57
	19 UΔU UGU AUA UAA 56
BVDV-2 850	44 AΔA UAU GGU GΔA 57
Singer	43 UΔU UGU AUA UAA 54
	16 UΔU UGU AAA UAA 57

 "pseudo stop-codon"

consensus sequence

U	U	A	A	U	A	U	U								
A	A	A	U		U	A		A	A						
G	G	G	G	G	G	U	G	G							
1	2	3	4	5	6	7	8	9	10	11	12				

Fig. 5E

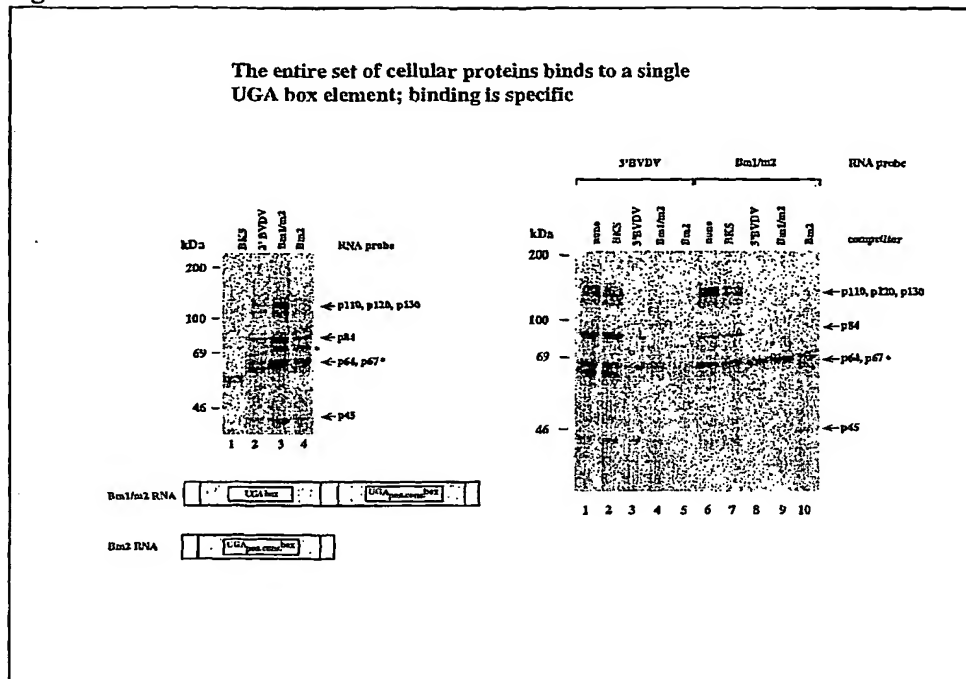


Fig. 6

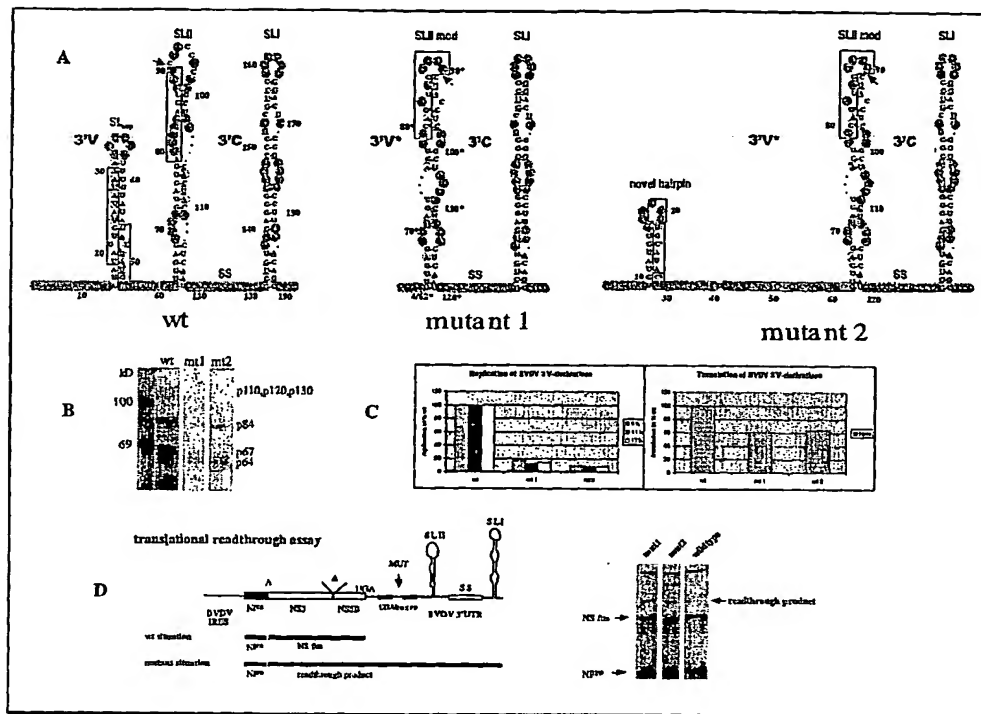


Fig. 8

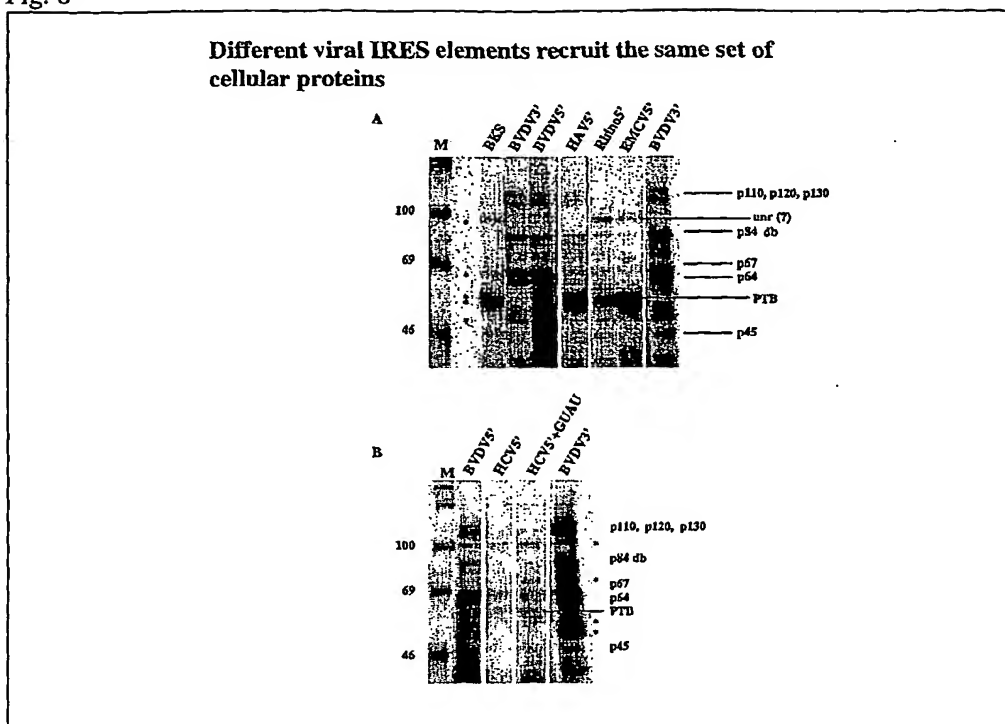


Fig. 9

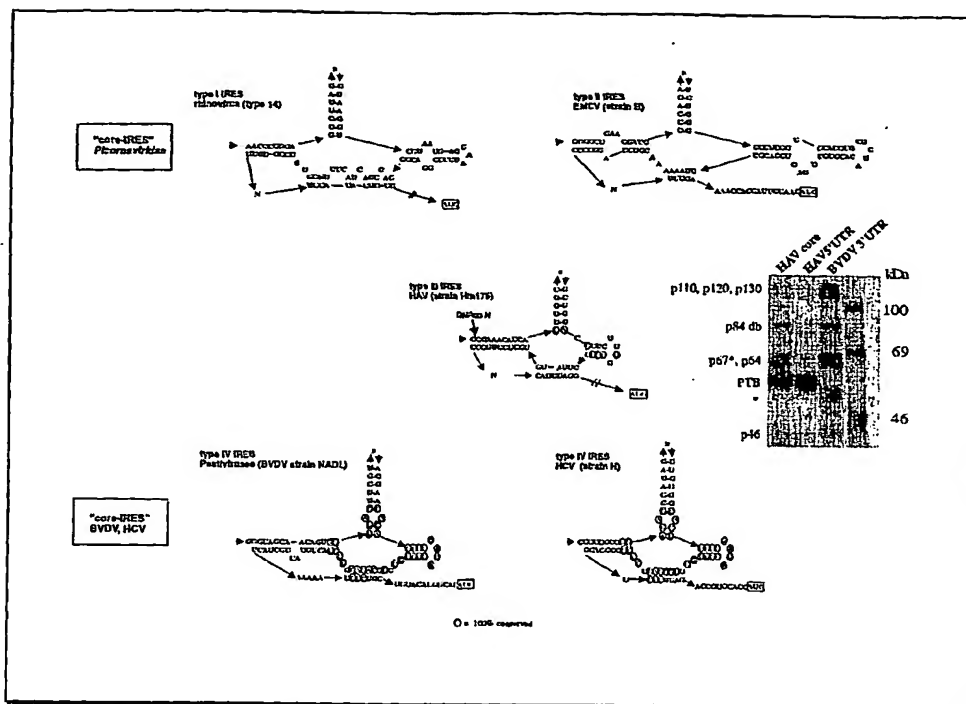


Fig. 10

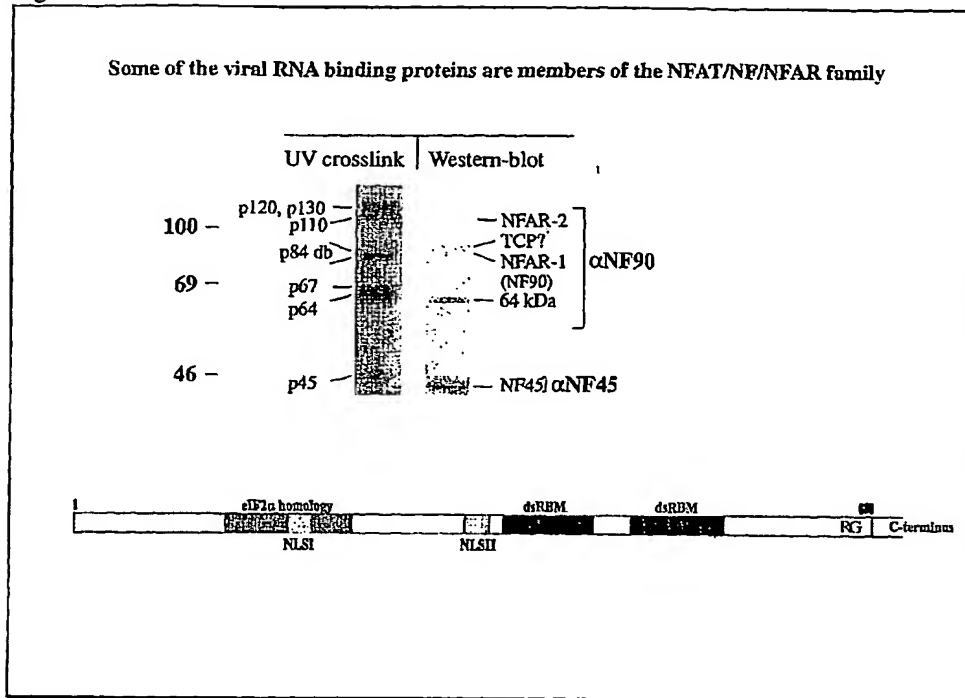


Fig. 10A

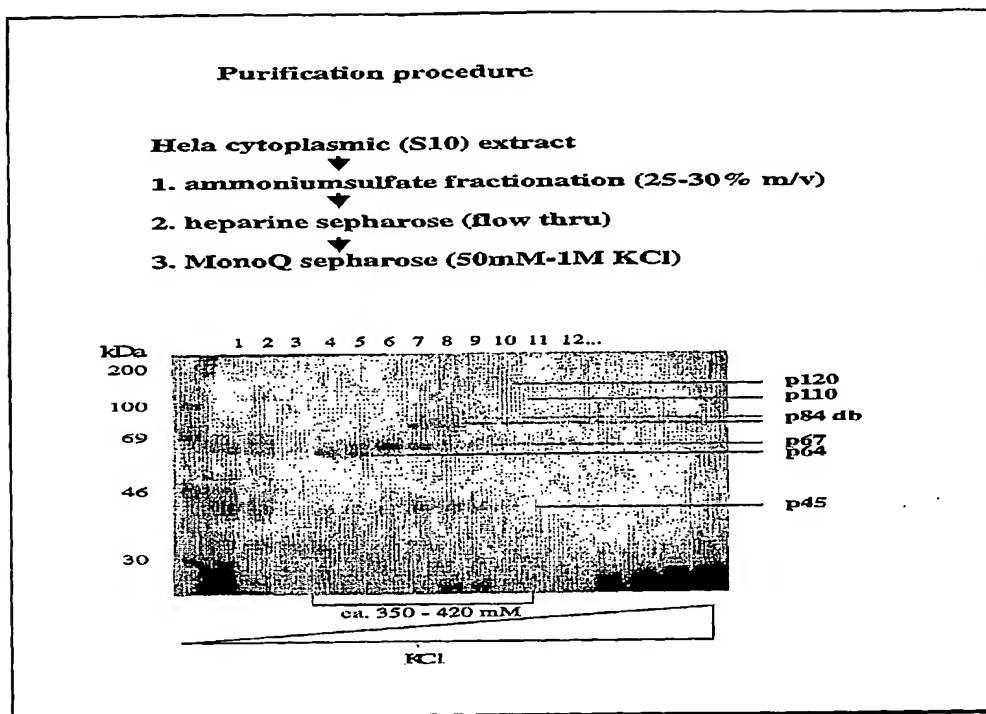


Fig. 11

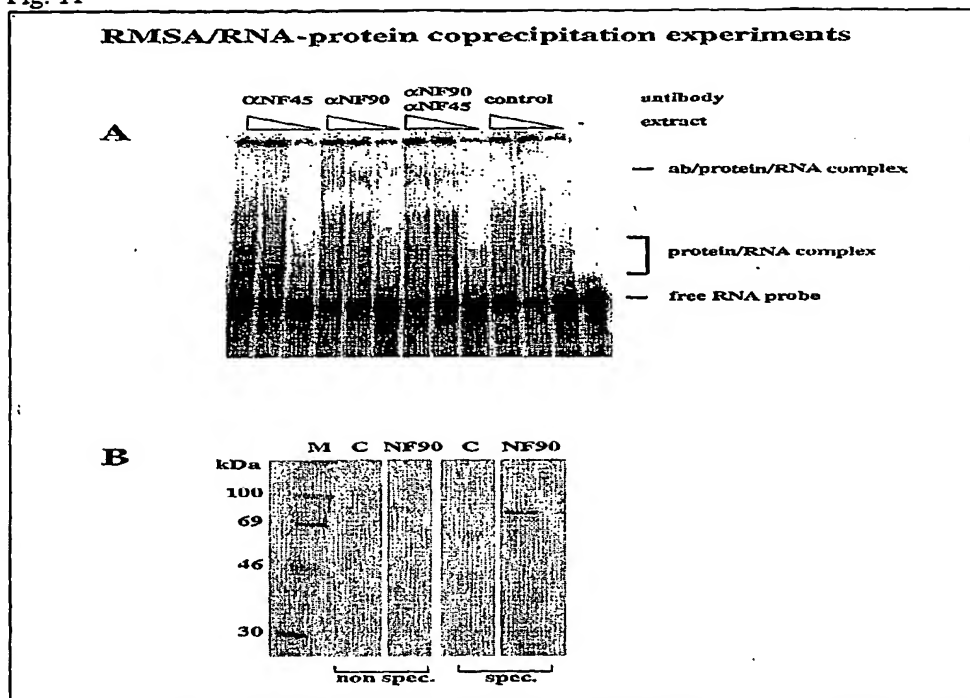


Fig. 12A

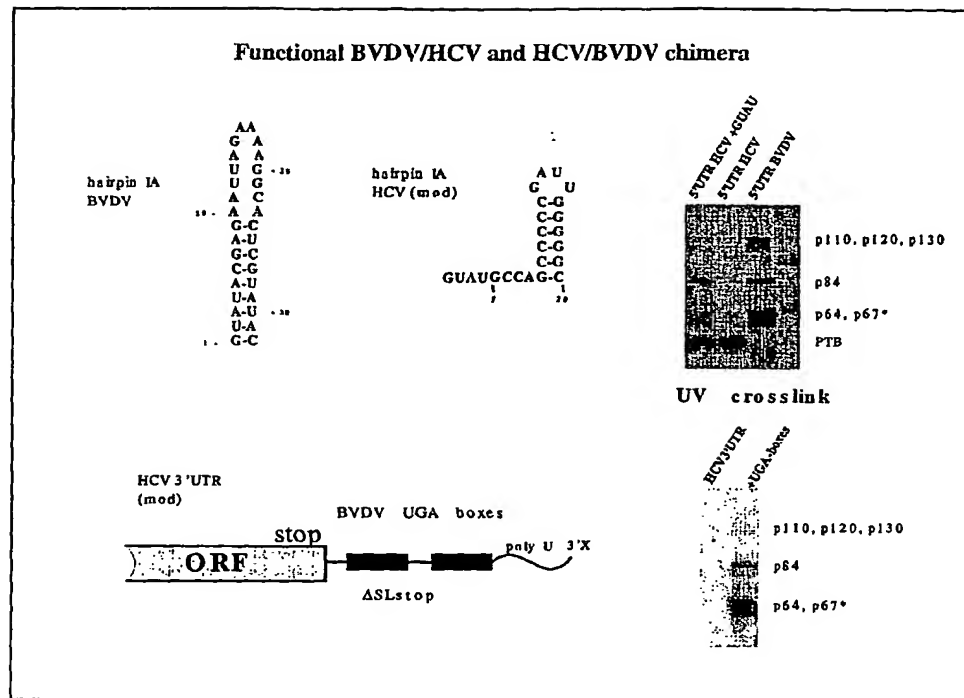


Fig. 12B

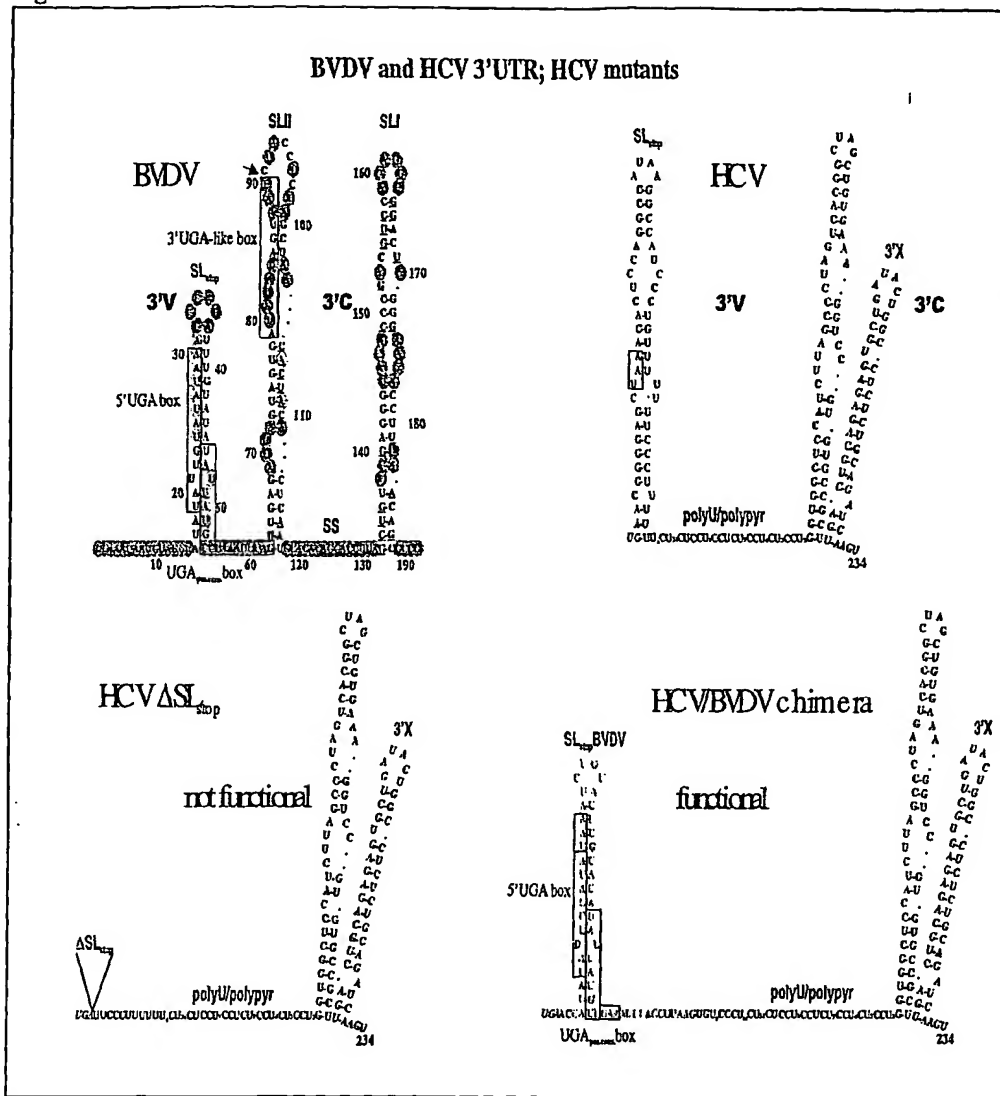
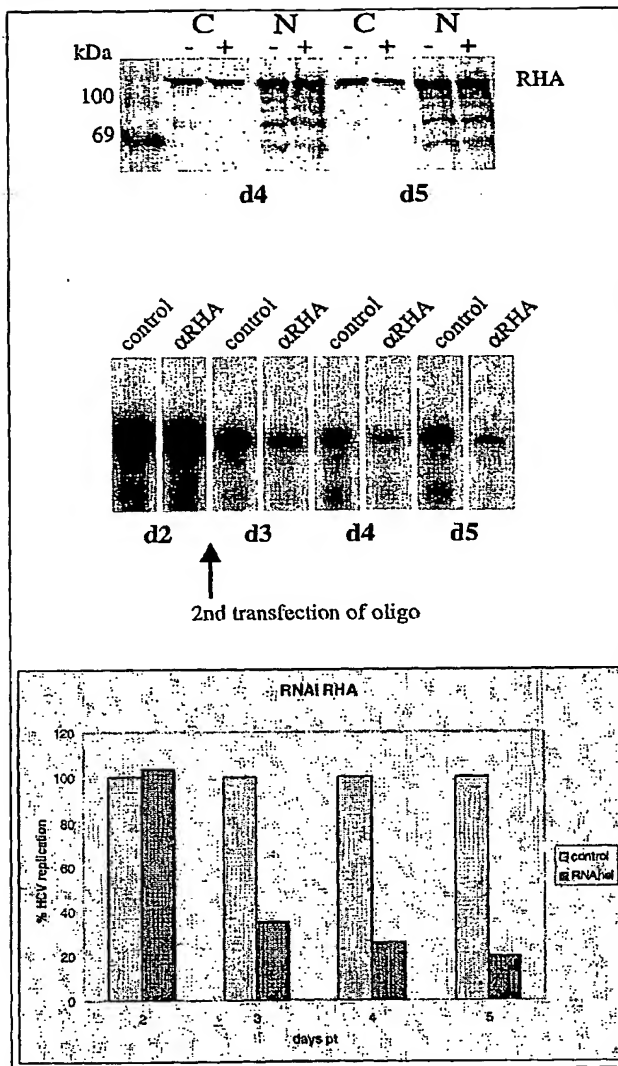


Fig. 13



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